FAX NO. :6502466597

Apr. 23 2004 11:58AM P2

Programs available for the BLAST search

The NCBI BLAST family of programs includes:

blastp

compares an amino acid query sequence against a protein sequence database

blastn

compares a nucleotide query sequence against a nucleotide sequence database

blastx

compares a nucleotide query sequence translated in all reading frames against a protein sequence database

tblastn

compares a protein query sequence against a nucleotide sequence database dynamically translated in all reading frames

tblastx

compares the six-frame translations of a nucleotide query sequence against the six-frame translations of a nucleotide sequence database. Please note that tblastx program cannot be used with the nr database on the BLAST Web page.

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FRUM . GTabexasi me

NCBL

BLAST 2 sequences

BLAST Entrez ?

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Matrix BLOSUM62 gap open: 11 gap extension: 1

x_dropoff: 50 expect: 10.000 wordsize: 3 Filter Align

Sequence 1 lcl|DEX0150_1

Length 1710

Sequence 2 lcl|US5733748 Fig1 Length 167

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Apr. 23 2004 11:59AM P4

NCBI BLAST 2 sequences BLAST Entrez ?

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Matrix BLOSUM62 gap open: 11 gap extension: 1

x_dropoff: 50 expect: 10.000 wordsize: 3 Filter Align

Sequence 1 lcl|DEX0150_1 Length 1710

Sequence 2 lcl|US5733748 Fig2 Length 235

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Apr. 23 2004 11:59AM P5

NCBI BLAST 2 sequences BLAST Entrez ?

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Matrix BLOSUM62 gap open: 11 gap extension: 1

x_dropoff: 50 expect: 10.000 wordsize: 3 Filter Align

Sequence 1 lcl|DEX0150_1 Length 1710

Sequence 2 lcl|US5733748 Fig4 Length 135

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Apr. 23 2004 11:59AM P6

NCBI BLAST 2 sequences BLAST Entrez ?

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Sequence 1 lcl|DEX0150_1 Length 1710

Sequence 2 lcl|US5733748 Fig5 Length 228

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NCB1

BLAST 2 sequences

BLAST Entrez ?

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Matrix BLOSUM62 ▼ gap open: 11 gap extension: 1

x_dropoff: 50 expect: 10.000 wordsize: 3 Filter Align

Sequence 1 lcl|DEX0150_1

Length 1710

Sequence 2 lel|US5733748 Fig6 Length 163

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Apr. 23 2004 11:59AM P8

NCBI

BLAST 2 sequences

BLAST Entre= ?

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Matrix BLOSUM62 gap open: 11 gap extension: 1

x_dropoff: 50 expect: 10.000 wordsize: 3 Filter Align

Sequence 1 lcI|DEX0150_1

Length 1710

Sequence 2 lol|US5733748 Fig8 Length 81

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Apr. 23 2004 11:59AM P9

NCBI

BLAST 2 sequences

BLAST Entrez ?

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Sequence 1 lcl|DEX0150_1

Length 1710

Sequence 2 lcl|US5733748 Fig9 Length 323

Ø 022

FROM :diaDexus, Inc.

FAX NO. :6502466597

Apr. 23 2004 12:00PM P10

NCBI

BLAST 2 sequences

BLAST Entrez

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Matrix BLOSUM62 gap open: 11 gap extension: 1

expect; 10.000 wordsize: 3 x_dropoff: 50 Filter □ Align

Sequence 1 lcl|DEX0150_1

Length 1710

Sequence 2 lcl|US5733748 Fig10 Length 201

LICATA & TYRRELL

FROM : diaDexus, Inc.

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Apr. 23 2004 12:00PM P11

FRUM :diabexus,ine

NCBI

BLAST 2 sequences

BLAST Entrez ?

BLAST 2 SEQUENCES RESULTS VERSION BLASTX 2.2.3 [Apr-24-2002]

Matrix BLOSUM62 ▼ gap open: 11 gap extension: 1

x_dropoff: 50 expect: 10.000 wordsize: 3 Filter Align

Sequence 1 lcl|DEX0150_1 Length 1710

Sequence 2 lcl|US5733748 Fig11 Length 156